

	Result No.	Score	Query	Match	Length	DB	ID	Description
-	1	666	100.0	318	21	AAY84558		A fragment of platelet-derived growth factor C (PDGF-C).
	2	666	100.0	339	21	AAB58438		Lung cancer associated protein.
	3	666	100.0	345	20	AAY33679		Human VEGF-E protein.
	4	666	100.0	345	20	AAY41766		Human PRO200 protein.
	5	666	100.0	345	20	AAY30023		Human vascular endothelial growth factor receptor.
	6	666	100.0	345	21	AAB48657		Human zvegf3, SEQ ID NO: 1.
	7	666	100.0	345	21	AAB24250		Human platelet-derived growth factor receptor.
	8	666	100.0	345	21	AAB44342		Human PRO200 (UNQ1).
	9	666	100.0	345	21	AAB10633		Human RAGE gene product.
	10	666	100.0	345	21	AAB10634		Human VEGF-X homolog.
	11	666	100.0	345	21	AAB10635		Human VEGF-X protein.

ALIGNMENTS

Title:	US-09-457-066-2_COPY_226_345	Scoring table:	BLOSUM62	Minimum DB seq length:	0
Perfect score:	666	Searched:	522463 seqs, 7407390 residues	Maximum DB seq length:	2000000000
Sequence:	1 AFVGRKSRRVVDNLNLTEEV DVALEHHEECDCVCRGSTGG 120	Total number of hits satisfying chosen parameters:	522463	Post-processing:	Minimum Match 0%
					Maximum Match 100%
					Listing first 45 summaries
Database :	A_Genesed_1101:*				
	1: /SIDS2/gcadata/geneseq/geneseq/AA1980.DAT:*				
	2: /SIDS2/gcadata/geneseq/geneseq/AA1981.DAT:*				
	3: /SIDS2/gcadata/geneseq/geneseq/AA1982.DAT:*				
	4: /SIDS2/gcadata/geneseq/geneseq/AA1983.DAT:*				
	5: /SIDS2/gcadata/geneseq/geneseq/AA1984.DAT:*				
	6: /SIDS2/gcadata/geneseq/geneseq/AA1985.DAT:*				
	7: /SIDS2/gcadata/geneseq/geneseq/AA1986.DAT:*				
	8: /SIDS2/gcadata/geneseq/geneseq/AA1987.DAT:*				
	9: /SIDS2/gcadata/geneseq/geneseq/AA1988.DAT:*				
	10: /SIDS2/gcadata/geneseq/geneseq/AA1989.DAT:*				
	11: /SIDS2/gcadata/geneseq/geneseq/AA1990.DAT:*				
	12: /SIDS2/gcadata/geneseq/geneseq/AA1991.DAT:*				
	13: /SIDS2/gcadata/geneseq/geneseq/AA1992.DAT:*				
	14: /SIDS2/gcadata/geneseq/geneseq/AA1993.DAT:*				
	15: /SIDS2/gcadata/geneseq/geneseq/AA1994.DAT:*				
	16: /SIDS2/gcadata/geneseq/geneseq/AA1995.DAT:*				
	17: /SIDS2/gcadata/geneseq/geneseq/AA1996.DAT:*				
	18: /SIDS2/gcadata/geneseq/geneseq/AA1997.DAT:*				
	19: /SIDS2/gcadata/geneseq/geneseq/AA1998.DAT:*				
	20: /SIDS2/gcadata/geneseq/geneseq/AA1999.DAT:*				
	21: /SIDS2/gcadata/geneseq/geneseq/AA2000.DAT:*				
	22: /SIDS2/gcadata/geneseq/geneseq/AA2001.DAT:*				

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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